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(FILE 'HOME' ENTERED AT 15:15:06 ON 17 JUL 2002)

FILE 'REGISTRY' ENTERED AT 15:15:36 ON 17 JUL 2002

L1 1 S 9030-45-9/RN

L2 1 S 3416-24-8/RN

L3 1 S 3616-42-0/RN

FILE 'HCAPLUS' ENTERED AT 15:19:08 ON 17 JUL 2002

FILE 'REGISTRY' ENTERED AT 15:19:32 ON 17 JUL 2002

L4 SET SMARTSELECT ON

SEL L1 1- CHEM : 17 TERMS

SET SMARTSELECT OFF

FILE 'HCAPLUS' ENTERED AT 15:19:33 ON 17 JUL 2002

L5 447 S L4

FILE 'REGISTRY' ENTERED AT 15:19:39 ON 17 JUL 2002

L6 SET SMARTSELECT ON

SEL L2 1- CHEM : 12 TERMS

SET SMARTSELECT OFF

FILE 'HCAPLUS' ENTERED AT 15:19:40 ON 17 JUL 2002

L7 19304 S L6

FILE 'REGISTRY' ENTERED AT 15:19:46 ON 17 JUL 2002

L8 SET SMARTSELECT ON

SEL L3 1- CHEM : 6 TERMS

SET SMARTSELECT OFF

FILE 'HCAPLUS' ENTERED AT 15:19:47 ON 17 JUL 2002

L9 620 S L8

L10 167 S L5 (L) L7 (L) L9

L11 119 S L10 AND PD<19970114

L12 605 S L7 (L) PREP/RL

L13 32 S L9 (L) PREP/RL

L14 134 S L5 (L) (L11 OR L12)

L15 17 S L5 (L) L11 (L) L12

L16 122 S L14 AND PD<19970114

L17 0 S L16 AND FERMENT?

L18 3 S L14 (L) FERMENT?

L19 190 S FERMENT? (L) (L7 OR L9)

L20 3 S L19 (L) L5

L21 122 S L16

L22 7 S L21 AND INHIBIT? AND PRODUCT

=> d' ibib ab 1-3

L20 ANSWER 1 OF 3 HCPLUS COPYRIGHT 2002 ACS  
ACCESSION NUMBER: 2001:634531 HCPLUS  
DOCUMENT NUMBER: 136:258038  
TITLE: Analysis of the chromosome sequence of the legume symbiont *Sinorhizobium meliloti* strain 1021  
AUTHOR(S): Capela, Delphine; Barloy-Hubler, Frederique; Gouzy, Jerome; Bothe, Gordana; Ampe, Frederic; Batut, Jacques; Boistard, Pierre; Becker, Anke; Boutry, Marc; Cadieu, Edouard; Dreano, Stephane; Gloux, Stephanie; Godrie, Therese; Goffeau, Andre; Kahn, Daniel; Kiss, Erno; Lelaure, Valerie; Masuy, David; Pohl, Thomas; Portetelle, Daniel; Puhler, Alfred; Purnelle, Benedicte; Ramsperger, Ulf; Renard, Clotilde; Thebault, Patricia; Vandebol, Micheline; Weidner, Stefan; Galibert, Francis  
CORPORATE SOURCE: Laboratoire de Biologie Moleculaire des Relations Plantes-Microorganismes, Unite Mixte de Recherche (UMR) 215 Centre National de la Recherche Scientifique (CNRS), Institut National de la Recherche Agronomique, Chemin, Tolosan, F-31326, Fr.  
SOURCE: Proceedings of the National Academy of Sciences of the United States of America (2001), 98(17), 9877-9882  
CODEN: PNASA6; ISSN: 0027-8424

PUBLISHER: National Academy of Sciences  
DOCUMENT TYPE: Journal  
LANGUAGE: English

AB *Sinorhizobium meliloti* is an alpha.-proteobacterium that forms agronomically important N<sub>2</sub>-fixing root nodules in legumes. We report here the complete sequence of the largest constituent of its genome, a 62.7% GC-rich 3654,135-bp circular chromosome. Annotation allowed assignment of a function to 59% of the 3341 predicted protein-coding ORFs, the rest exhibiting partial, weak, or no similarity with any known sequence. Unexpectedly, the level of reiteration within this replicon is low, with only two genes duplicated with more than 90% nucleotide sequence identity, transposon elements accounting for 2.2% of the sequence, and a few hundred short repeated palindromic motifs (RIME1, RIME2, and C) widespread over the chromosome. Three regions with a significantly lower GC content are most likely of external origin. Detailed annotation revealed that this replicon contains all housekeeping genes except two essential genes that are located on pSymB. Amino acid/peptide transport and degrdn. and sugar metab. appear as two major features of the *S. meliloti* chromosome. The presence in this replicon of a large no. of nucleotide cyclases with a peculiar structure, as well as of genes homologous to virulence determinants of animal and plant pathogens, opens perspectives in the study of this bacterium both as a free-living soil microorganism and as a plant symbiont.

REFERENCE COUNT: 53 THERE ARE 53 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L20 ANSWER 2 OF 3 HCPLUS COPYRIGHT 2002 ACS  
ACCESSION NUMBER: 2000:68590 HCPLUS  
DOCUMENT NUMBER: 132:121532  
TITLE: Glucosamine fermentation with recombinant microorganisms with mutations in the glucosamine-6-phosphate metabolic pathway  
INVENTOR(S): Berry, Alan; Burlingame, Richard P.; Millis, James R.  
PATENT ASSIGNEE(S): DCV, Inc. D/B/A Bio-Technical Resources, USA  
SOURCE: PCT Int. Appl., 151 pp.  
CODEN: PIXXD2  
DOCUMENT TYPE: Patent  
LANGUAGE: English  
FAMILY ACC. NUM. COUNT: 3  
PATENT INFORMATION:

| PATENT NO. | KIND | DATE | APPLICATION NO. | DATE |
|------------|------|------|-----------------|------|
|------------|------|------|-----------------|------|

|  |    |          |                 |             |
|--|----|----------|-----------------|-------------|
| WO 2000004182  | A1 | 20000127 | WO 1999-US15976 | 19990715    |
| W: AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ,<br>DE, DK, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS,<br>JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK,<br>MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ,<br>TM, TR, TT, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD,<br>RU, TJ, TM |    |          |                 |             |
| RW: GH, GM, KE, LS, MW, SD, SL, SZ, UG, ZW, AT, BE, CH, CY, DE, DK,<br>ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG,<br>CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG   |    |          |                 |             |
| US 6372457   | B1 | 20020416 | US 1998-115475  | 19980715    |
| AU 9951028   | A1 | 20000207 | AU 1999-51028   | 19990715    |
| EP 1095158   | A1 | 20010502 | EP 1999-935577  | 19990715    |
| R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,<br>IE, SI, LT, LV, FI, RO   |    |          |                 |             |
| JP 2002520067  | T2 | 20020709 | JP 2000-560279  | 19990715    |
| PRIORITY APPLN. INFO.:   |    |          | US 1998-115475  | A 19980715  |
|  |    |          | US 1997-35494P  | P 19970114  |
|  |    |          | WO 1998-US800   | A2 19980114 |
|  |    |          | WO 1999-US15976 | W 19990715  |

AB The present invention relates to a method and materials for producing **glucosamine** by **fermn.** of a genetically modified microorganism. Included in the present invention are genetically modified microorganisms useful in the present method for producing **glucosamine**, as well as recombinant nucleic acid mols. and the proteins produced by such recombinant nucleic acid mols. Thus, a modified Escherichia coli strain with the nagA-D genes deleted, the manXYZ genes mutationally inactivated, and the glmS gene replaced with an inducible mutant glmS gene encoding a **glucosamine-6-phosphate synthase** resistant to **glucosamine-6-phosphate** inhibition was constructed. In **fermentor** cultures, **glucosamine** concns. in excess of 12 g/L were obtained with this E. coli strain.

REFERENCE COUNT: 5 THERE ARE 5 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L20 ANSWER 3 OF 3 HCPLUS COPYRIGHT 2002 ACS  
 ACCESSION NUMBER: 1998:493700 HCPLUS  
 DOCUMENT NUMBER: 129:121714  
 TITLE: Process for production of N-glucosamine  
 INVENTOR(S): Berry, Alan; Burlingame, Richard P.; Millis, James R.  
 PATENT ASSIGNEE(S): Bio-Technical Resources, USA; Berry, Alan; Burlingame, Richard P.; Millis, James R.  
 SOURCE: PCT Int. Appl., 91 pp.  
 CODEN: PIXXD2  
 DOCUMENT TYPE: Patent  
 LANGUAGE: English  
 FAMILY ACC. NUM. COUNT: 3  
 PATENT INFORMATION:

| PATENT NO.  | KIND | DATE     | APPLICATION NO. | DATE       |
|---|------|----------|-----------------|------------|
| WO 9830713  | A1   | 19980716 | WO 1998-US800   | 19980114   |
| W: AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE,<br>DK, EE, ES, FI, GB, GE, GH, GM, GW, HU, ID, IL, IS, JP, KE, KG,<br>KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX,<br>NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT,<br>UA, UG, US, UZ, VN, YU, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM |      |          |                 |            |
| RW: GH, GM, KE, LS, MW, SD, SZ, UG, ZW, AT, BE, CH, DE, DK, ES, FI,<br>FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM,<br>GA, GN, ML, MR, NE, SN, TD, TG  |      |          |                 |            |
| AU 9859604  | A1   | 19980803 | AU 1998-59604   | 19980114   |
| US 6372457  | B1   | 20020416 | US 1998-115475  | 19980715   |
| PRIORITY APPLN. INFO.:  |      |          | US 1997-35494P  | P 19970114 |
|   |      |          | WO 1998-US800   | W 19980114 |

AB The present invention relates to a method for producing glucosamine by

fermn. of a genetically modified microorganism.

L22 ANSWER 1 OF 7 HCPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1997:392564 HCPLUS

DOCUMENT NUMBER: 127:47021

TITLE: Substrate binding is required for assembly of the active conformation of the catalytic site in Ntn amidotransferases: evidence from the 1.8 .ANG. crystal structure of the glutaminase domain of **glucosamine 6-phosphate synthase**

[Erratum to document cited in CA125:136326]

AUTHOR(S): Isupov, Michail N.; Obmolova, Galya; Butterworth, Susanna; Badet-Denisot, Marie-Ange; Badet, Bernard; Polikarpov, Igor; Littlechild, Jennifer A.; Teplyakov, Alexei

CORPORATE SOURCE: Dep. Chem. Biological Scis., Univ. Exeter, Exeter, EX4 4QD, UK

SOURCE: Structure (London) (1997), 5(5), 723  
CODEN: STRUE6; ISSN: 0969-2126

PUBLISHER: Current Biology

DOCUMENT TYPE: Journal

LANGUAGE: English

AB The catalytic mechanism described for **glucosamine 6-phosphate synthase** was based on the mechanism of penicillin hydrolysis by penicillin acylase proposed by Duggleby et al. (1995) to which ref. should have been made: Duggleby, H.J., Tolley, S.P., Hill, C.P., Dodson, E.J., Dodson, G. and Moody, P.C.E. (1995) Nature 373, 264-268.

L22 ANSWER 2 OF 7 HCPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1996:460296 HCPLUS

DOCUMENT NUMBER: 125:136326

TITLE: Substrate binding is required for assembly of the active conformation of the catalytic site in Ntn amidotransferases: evidence from the 1.8 .ANG. crystal structure of the glutaminase domain of **glucosamine 6-phosphate synthase**

AUTHOR(S): Isupov, Michail N.; Obmolova, Gayla; Butterworth, Susanna; Badet-Denisot, Marie-Ange; Badet, Bernard; Polikarpov, Igor; Littlechild, Jennifer A.; Teplyakov, Alexei

CORPORATE SOURCE: Dep. Chem. Biological Scis., Univ. Exeter, Exeter, EX4 4QD, UK

SOURCE: Structure (London) (1996), 4(7), 801-810  
CODEN: STRUE6; ISSN: 0969-2126

DOCUMENT TYPE: Journal

LANGUAGE: English

AB Amidotransferases use the amide nitrogen of glutamine in a no. of important biosynthetic reactions. They are composed of a glutaminase domain, which catalyzes the hydrolysis of glutamine to glutamate and ammonia, and a synthetase domain, catalyzing amination of the substrate. To gain insight into the mechanism of nitrogen transfer, we examd. the structure of the glutaminase domain of **glucosamine 6-phosphate synthase** (GLMS). The crystal structures of the enzyme complexed with glutamate and with a competitive inhibitor, Glu-hydroxamate, have been detd. to 1.8 .ANG. resoln. The protein fold has structural homol. to other members of the superfamily of N-terminal nucleophile (Ntn) hydrolases, being a sandwich of antiparallel .beta. sheets surrounded by two layers of .alpha. helices. The structural homol. between the glutaminase domain of GLMS and that of phosphoribosyl pyrophosphate (PRPP) amidotransferase (the only other Ntn amidotransferase whose structure is known) indicates that they may have diverged from a common ancestor. Cys1 is the catalytic nucleophile in GLMS, and the nucleophilic character of its thiol group appears to be increased through general base activation by its own .alpha.-amino group.

Cys1 can adopt two conformations, one active and one inactive; glutamine binding locks the residue in a predetd. conformation. We propose that when a nitrogen acceptor is present Cys1 is kept in the active conformation, explaining the phenomenon of substrate-induced activation of the enzyme, and that Arg26 is central in this coupling.

L22 ANSWER 3 OF 7 HCPLUS COPYRIGHT 2002 ACS  
ACCESSION NUMBER: 1994:1295 HCPLUS  
DOCUMENT NUMBER: 120:1295  
TITLE: Glucose regulation of transforming growth factor-.alpha. expression is mediated by products of the hexosamine biosynthesis pathway  
AUTHOR(S): Daniels, Marc C.; Kansal, Preeti; Smith, Tom M.; Paterson, Andrew J.; Kudlow, Jeffrey E.; McClain, Donald A.  
CORPORATE SOURCE: Veterans Adm. Med. Cent., Birmingham, AL, 35294, USA  
SOURCE: Mol. Endocrinol. (1993), 7(8), 1041-8  
CODEN: MOENEN; ISSN: 0888-8809  
DOCUMENT TYPE: Journal  
LANGUAGE: English  
AB The authors have recently shown that glucose and **glucosamine** regulate the transcription of transforming growth factor-.alpha. (TGF.alpha.) in rat aortic smooth muscle (RASM) cells. Based on the increased potency of **glucosamine** compared to glucose, the authors hypothesized that stimulation of TGF.alpha. transcription by glucose is mediated through the hexosamine biosynthesis pathway. The yeast cDNA for the rate-limiting enzyme of this pathway, **glutamine :fructose-6-phosphate amidotransferase** (GFA), was therefore expressed in RASM cells. GFA-transfected cells showed an increase in GFA activity, exhibiting a 2.2-fold increase in the synthesis of **glucosamine-6-phosphate**, the first **product** of the hexosamine biosynthetic pathway. To test the effect of GFA overexpression on TGF.alpha. transcriptional activity, cells were transiently cotransfected with GFA along with a reporter plasmid contg. the firefly luciferase gene under control of the TGF.alpha. promoter. GFA-transfected cells exhibited a glucose-dependent 2-fold increase in TGF.alpha. activity compared to control cells. Maximal stimulation of TGF.alpha. luciferase activity by **glucosamine**, however, was equiv. in GFA- and control-transfected cells, confirming that the stimulation obsd. by both agents operated through the same pathway. This increase in TGF.alpha. activity was inhibited (85% at 0.5 mM glucose and 69% at 30 mM glucose) by the glutamine analog and **inhibitor** of GFA, 6-diazo-5-oxonorleucine (10 .mu.M). Control studies confirmed that the increased TGF.alpha.-luciferase activity in the GFA-expressing cells was not an artifact of altered growth, survival, or transfection efficiency. Expts. using pharmacol. agents to stimulate or **inhibit** protein kinase C and cAMP-dependent kinase do not support a role for these second messengers in the signaling pathway. Tunicamycin inhibited the ability of glucose to stimulate TGF.alpha. activity, suggesting that protein glycosylation does play a role. The authors conclude that **products** of the hexosamine biosynthesis pathway mediate the stimulation by glucose of TGF.alpha. in aortic smooth muscle cells.

L22 ANSWER 4 OF 7 HCPLUS COPYRIGHT 2002 ACS  
ACCESSION NUMBER: 1993:119586 HCPLUS  
DOCUMENT NUMBER: 118:119586  
TITLE: Investigation of the **inhibition** pathway of **glucosamine synthase** by N3-(4-methoxyfumaroyl)-L-2,3-diaminopropanoic acid by semiempirical quantum mechanical and molecular mechanics methods  
AUTHOR(S): Tarnowska, M.; Oldziej, S.; Liwo, A.; Grzonka, Z.; Borowski, E.  
CORPORATE SOURCE: Dep. Chem., Univ. Gdansk, Gdansk, PL-80-952, Pol.  
SOURCE: Eur. Biophys. J. (1992), 21(4), 273-80

CODEN: EBJOE8; ISSN: 0175-7571

DOCUMENT TYPE:

Journal

LANGUAGE:

English

AB **Glucosamine 6-phosphate synthase**

(EC 2.6.1.16) is a promising target in antifungal drug design. It has been reported that its potent **inhibitor**, N3-(4-methoxyfumaroyl)-L-2,3-diaminopropionic acid (FMDP), inactivates the enzyme by the Michael addn. of the SH group to the FMDP mol. followed by cyclization reactions. Here, using semiempirical MNDO, PM3, and mol. mechanics methods, the energetics and kinetic possibility of the formation of various stereoisomers of the **products** of cyclization of the Michael addn. **products** detected exptl. were investigated. It was found that the substituted 1,4-thiazin-3-one can be formed in 1 step under alk. conditions; the stereoisomers of this compd., predicted to be the most stable on the basis of theor. calcns., were also the dominant ones in reality.

L22 ANSWER 5 OF 7 HCPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1990:194297 HCPLUS

DOCUMENT NUMBER: 112:194297

TITLE: **Glucosamine-6-phosphate**

**synthase** from *Escherichia coli*: determination of the mechanism of inactivation by N3-fumaroyl-L-2,3-diaminopropionic derivatives

AUTHOR(S): Kucharczyk, Nathalie; Denisot, Marie Ange; Le Goffic, Francois; Badet, Bernard

CORPORATE SOURCE: Lab. Bioorg. Biotechnol., ENSCP, Paris, 75231, Fr.

SOURCE: Biochemistry (1990), 29(15), 3668-76

CODEN: BICHAW; ISSN: 0006-2960

DOCUMENT TYPE: Journal

LANGUAGE: English

AB A mechanistic investigation of the inactivation of *E. coli*

**glucosamine-6-phosphate synthase** by

N3-(4-methoxyfumaroyl)-L-2,3-diaminopropionate (FMDP) was undertaken. On the basis of the known participation of the N-terminal cysteine residue in this process, model reactions between FMDP and L-cysteine and between FMDP and the synthetic decapeptide, Cys-Gly-Ile-val-Gly-Ala-Ile-Ala-Gln-Arg, corresponding to the N-terminal protein sequence, were studied. The results allowed a pathway to be proposed that was in perfect agreement with the biochem. results: enzyme inactivation arose from Michael addn. of glutamine-binding site cysteine-1 on the fumaroyl double bond at the .beta.-position of the ester group. Upon denaturation under slightly alk. conditions, this adduct underwent cyclization to a transient succinimide adduct, which rearranged into the stable 2-substituted 1,4-thiazin-3-one-5-carboxylate involving participation of the cysteine amino group. The tryptic radiolabeled peptides purified from [3H]FMDP-treated enzyme and resistant to Edman degrdn. coeluted with the **products** resulting from the model reaction between the synthetic decapeptide and the **inhibitor**.

L22 ANSWER 6 OF 7 HCPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1988:146073 HCPLUS

DOCUMENT NUMBER: 108:146073

TITLE: **Glucosamine synthetase** from

*Escherichia coli*: kinetic mechanism and **inhibition** by N3-fumaroyl-L-2,3-diaminopropionic derivatives

AUTHOR(S): Badet, Bernard; Vermoote, Patricia; Le Goffic, Francois

CORPORATE SOURCE: Lab. Bioorg. Biotechnol., ENSCP, Paris, 75231, Fr.

SOURCE: Biochemistry (1988), 27(7), 2282-7

CODEN: BICHAW; ISSN: 0006-2960

DOCUMENT TYPE: Journal

LANGUAGE: English

AB N3-(4-Methoxyfumaroyl)-L-2,3-diaminopropionic acid (FMDP), a member of a new class of glutamine analogs, was investigated as an **inhibitor** of pure *E. coli* glucosamine phosphate synthetase (I). **Product**

and dead-end **inhibition** studies indicated an ordered assocn. to the enzyme with the sugar mol. binding prior to substrate or **inhibitor**. The inactivation exhibited pseudo-1st-order kinetics, was irreversible, and occurred faster in the presence of fructose 6-phosphate, a behavior previously reported for the partially purified enzyme from *Salmonella typhimurium*. FMDP was found to be one of the most efficient **inhibitors** of I to date. The **inhibition** occurred with partial covalent incorporation of L-FMDP into I. In the presence of fructose 6-phosphate, enzyme inactivation with [2-3H]-DL-FMDP was assocd. with the incorporation of 0.75 equiv of **inhibitor** and with the modification of 0.78 SH residue per enzyme subunit. This result is the 1st evidence for covalent entrapment of the entire **inhibitor** mol. following FMDP-mediated I inactivation. Preliminary inactivation with 6-diazo-5-oxo-L-norleucine, known to alkylate selectively the N-terminal cysteine residue, completely prevented radioactivity incorporation. Therefore, this **inhibitor** is postulated to covalently modify I through direct addn. of the thiol nucleophile from the terminal cysteine residue to the Michaelis acceptor, so acting as an affinity label rather than a mechanism-based **inhibitor**.

L22 ANSWER 7 OF 7 HCAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1985:204266 HCAPLUS  
DOCUMENT NUMBER: 102:204266  
TITLE: Synthesis of 3,4-iminocyclohexyl-glycine and its N-benzyloxycarbonyl derivative  
AUTHOR(S): Dzieduszycka, Maria; Martelli, Sante; Borowski, Edward  
CORPORATE SOURCE: Dep. Pharm. Technol. Biochem., Tech. Univ. Gdansk,  
Gdansk, Pol.  
SOURCE: Int. J. Pept. Protein Res. (1985), 25(1),  
99-104  
CODEN: IJPPC3; ISSN: 0367-8377  
DOCUMENT TYPE: Journal  
LANGUAGE: English  
OTHER SOURCE(S): CASREACT 102:204266  
AB The title compds. I [R = H, PhCH<sub>2</sub>O<sub>2</sub>C (Z)] were prep'd. from prep'd. from cyclohexenylglycines II (R<sub>1</sub> = Z, CF<sub>3</sub>CO) via an addn. reaction with iodine isocyanate (III). Thus, III was added to II (R<sub>1</sub> = Z) to give addn. products IV (R<sub>2</sub> = Z, R<sub>3</sub> = NCO) as a mixt. of the 2 possible 3- and 4-positional isomers. The latter were treated with MeOH to give the corresponding IV (R<sub>2</sub> = Z, R<sub>3</sub> = NHCO<sub>2</sub>Me) (as 2 isomers), which were cyclized in the presence of KOH to give I (R = Z). II (R<sub>1</sub> = CF<sub>3</sub>CO) was converted to I (R = H) via IV (R<sub>2</sub> = CF<sub>3</sub>CO, R<sub>3</sub> = NHCO<sub>2</sub>Me). I (R = H) inhibited glucosamine synthetase.